

SEQUENCE LISTING

5 (1) GENERAL INFORMATION:

- (i) APPLICANT: HARADA, SHUN-ICHI
SAMPATH, K. T.
RODAN, GIDEON A.
- 10 (ii) TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR IDENTIFYING
MORPHOGEN ANALOGS
- (iii) NUMBER OF SEQUENCES: 8
- 15 (iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES
(B) STREET: 45 SOUTH STREET
(C) CITY: HOPKINTON
(D) STATE: MA
20 (E) COUNTRY: USA
(F) ZIP: 01748
- (v) COMPUTER READABLE FORM:
25 (A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- 30 (vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:
- 35 (viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: VITO, CHRISTINE C.
(B) REGISTRATION NUMBER: 39,061
(C) REFERENCE/DOCKET NUMBER: CRP-126
- 40 (ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: (617)-248-7000
(B) TELEFAX: (617)-248-7100

45 (2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 1067 base pairs
(B) TYPE: nucleic acid
50 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- 55 (ix) FEATURE:
(A) NAME/KEY: misc_feature
(B) LOCATION: 1..1067
(D) OTHER INFORMATION: /product= "MOUSE TYPE 10 COLLAGEN
60 PROMOTER"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

65	TCGATCCTAA AACACTTAAG GATATTCTG TAAGGCTGTG AAAGAGAAAA CCAACTACTT	60
	ACACGGATGG AGACCATGTT TATTCTTTG GGAGAAAAGC CTAATTGGGA CGCTTCGAGA	120
	TCCCTATAGG AAATGCACC AGTAGTCAAC TGGATTTTAA AAAGGCAAAG CTTGAGGATT	180
70	TTTTTTTCCC TTTGAAATGA ATGTAGCAAA CTTATGTAAG CACGGAATAG GATTATTAGT	240

5 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 11 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

10 (ix) FEATURE:
(A) NAME/KEY: misc_feature
(B) LOCATION: 1..11
OLIGO" (D) OTHER INFORMATION: /product= "MEF-2 CONSENSUS MUTANT

15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
TTAAACATAA A

11

20 (2) INFORMATION FOR SEQ ID NO:8:

25 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 11 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

30 (ix) FEATURE:
(A) NAME/KEY: misc_feature
(B) LOCATION: 1..11
... (D) OTHER INFORMATION: /product= "MEF-2 MUTANT CONSENSUS"

35 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
CTAAACATAA C

11